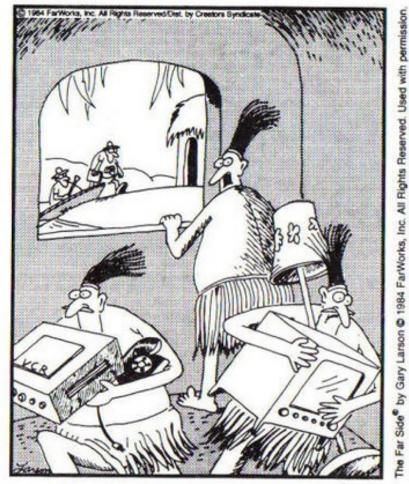
THE FAR SIDE" BY GARY LARSON

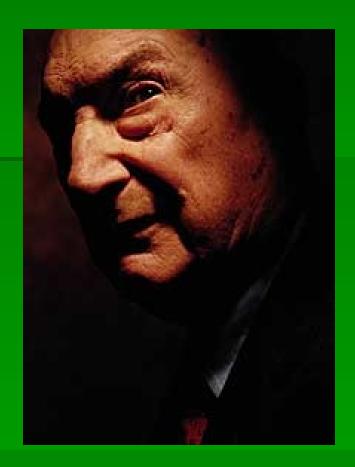


"Anthropologists! Anthropologists!"

Let food be your medicine and medicine be your food.

- Hippocrates, 400 BCE

"One future intelligence problem: knowing what drugs the other guys are on."



SAME SPECIES...VERY DIFFERENT CREATURES

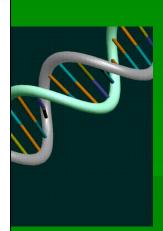
Nurture



Nature







Assumptions

- Common dietary compounds act on the human genome, directly or indirectly, to alter gene expression or structure;
- Some individuals, under some circumstances, can have diet become a serious risk factor;
- Some diet-regulated genes and their normal common variants help shape processes like susceptibility to disease/injury/extreme environments and progression, recovery from, and severity of breakdown
- The degree to which diet influences these processes depends in part on an individual's genetic makeup
- Dietary interventions based on the "nutriome" can be used to prevent, mitigate, or cure disease/injury¹
 - Kaput and Rodriguez, 2004 "Nutritional Genomics", *Physiol Genomics* 16(2):166-167

What this briefing is NOT about

Thorough literature review

GKMM Ncol

PPARά G/C

ACE I/D

ADRA2A

ACTN3 R577X NO Glu289ASp ADRB2

APOE

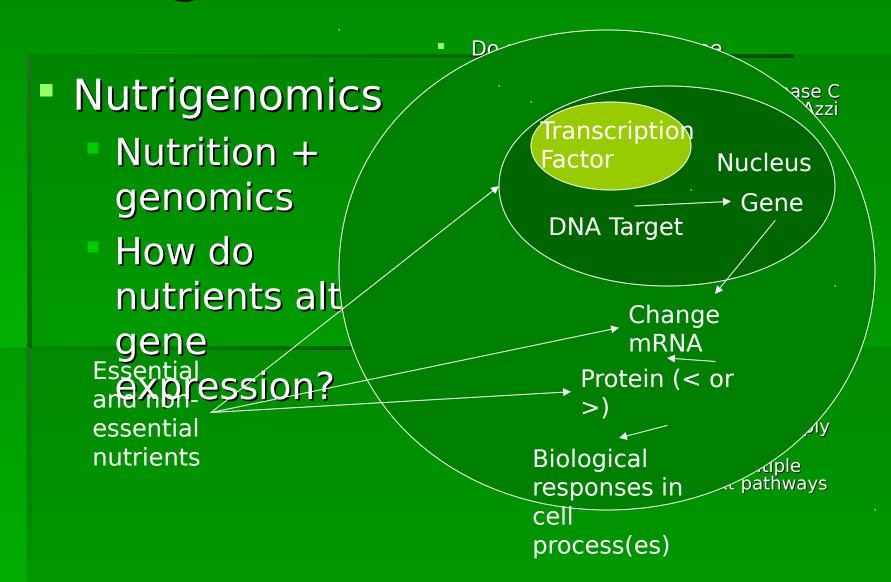
Spl transc

List of Single

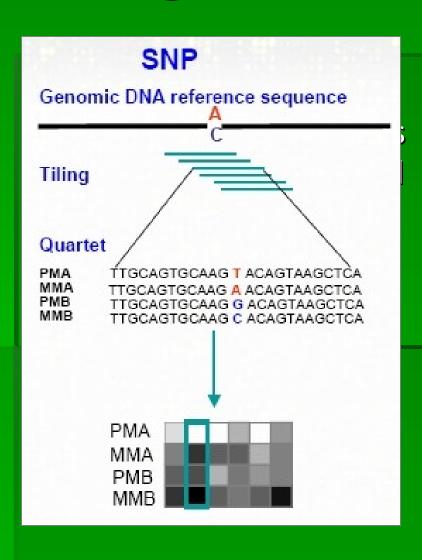
1: Steiner C, Amould S, Scalbert A, Manach C Isoflavones and the prevention of breast and prostate cancer: new perspectives opened by nutrigenomics. Br J Nutr. 2008 May;99 E Suppl 1:ES78-108. PMID: 18503737 [PubMed - in process] 2: Ovesná J, Slabý O, Toussaint O, Kodícek M, Marsík P, Pouchová V, Vaněk T. High throughput 'omics' approaches to assess the effects of phytochemicals in human health studies. Br J Nutr. 2008 May;99 E Suppl 1:ES127-34. PMID: 18503733 [PubMed - in process] Mortensen A, Sorensen IK, Wilde C, Dragoni S, Mullerová D, Toussaint O, Zloch Z, Sgaragli G, Ovesná J. Biological models for phytochemical research: from cell to human organism. Br J Nutr. 2008 May;99 E Suppl 1:ES118-26 PMID: 18503732 [PubMed - in process] 4: Stienstra R, Duval C, Keshtkar S, van der Laak J, Kersten S, Müller M. PPARgamma activation promotes infiltration of alternatively activated macrophages into adipose tissue. J Biol Chem. 2008 Jun 9. [Epub ahead of print] PMID: 18541527 [PubMed - as supplied by publisher] 14: Sát Lez J, Cabrer JM, Rosselló CA, Palou A, Picó C. Formation of him agroban indiffucts of activities in the its ingestion in rats is depinding on age and sex J Agric Food Chen., 2019 Jul 9;56(13):1096-111. Ep.: 2008 Jun 10 PMID: 18540624 [PubMed - in process] Omega-3: from cod-liver oil to nutrigenomics. Minerva Pediatr. 2008 Aug;60(4):443-455. PMID: 18511896 [PubMed - as supplied by publisher] 7: Martinez JA, Parra MD, S n os J , Morer o-Aliaga MJ, Marti A, Martinez-Gonzalez MA Genotype-dependent response to energy-restricted diets in obese subjects: towards personalized nutrition Asia Pac J Clin Nutr. 2008;17 Suppl 1:119-22. Review. PMID: 18296317 [PubMed - indexed for MEDLINE] 8: Lee YY, Tsou CS, Lin HC, Ien CH, Wu YT. Global perspective of health related edible plants from the agricultural point of view. Asia Pac J Clin Nutr. 2008;17 Suppl 1:95-8. Review. PMID: 18296311 [PubMed - indexed for MEDLINE] Nutrigunonics the cutting edge and Alian perspectives.

Asia Pac J Clin Nutr. 2008;17 Suppl 1:12-5. Renaw. PMID: 18296291 [PubMed - indexed for MEDI INE]

Nutrigenomics (Nurture)



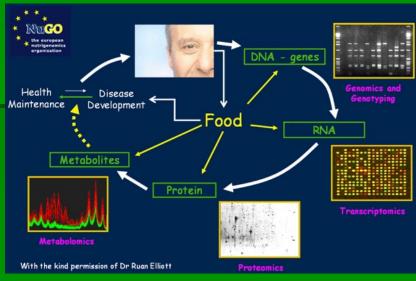
Nutrigenetics (Nature)



- SNPs occur once every 1k to 2k nucleotides, but occur at a frequency > 1% in the population
- Effects can be variable and not always dramatic
 - Can alter protein structure and function when the nucleotide base substitution occurs in a gene's coding region
 - When substitution occurs as part of the gene's regulatory promoter, the SNP may affect the conditions under which the protein is made

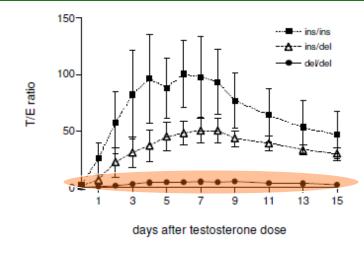
Omics: you can't stop with just one

- Nutrigenomics
- Epigenomics
- Nutritional epigenomics (as they influence DNA methylation, histone modification, and RNAassociated silencing)
- Transcriptomics
- Proteomics
- Metabolomics
- Microbiomics
- Connectomics
- HPomics



So do genes matter?





1977 1997

Examples of Type 1, 2, and 3 subjects in response to cognitive testing after multiple nights of sleep restriction. Van Dongen, H.

P. A., Maislin, G., Mullington, J. M., & Dinges, D. F. (2003). The cumulative cost of additional wakefulness: Dose-response effects on neurobehavioral functions and sleep physiology from chronic sleep restriction and total sleep deprivation. *SLEEP*, 26, 117-126.

- T:E ratio
- 55 healthy male volunteers
- Ins/ins (two copies of

2007 AU G. 2815 70 Gene) V. Del/del (no

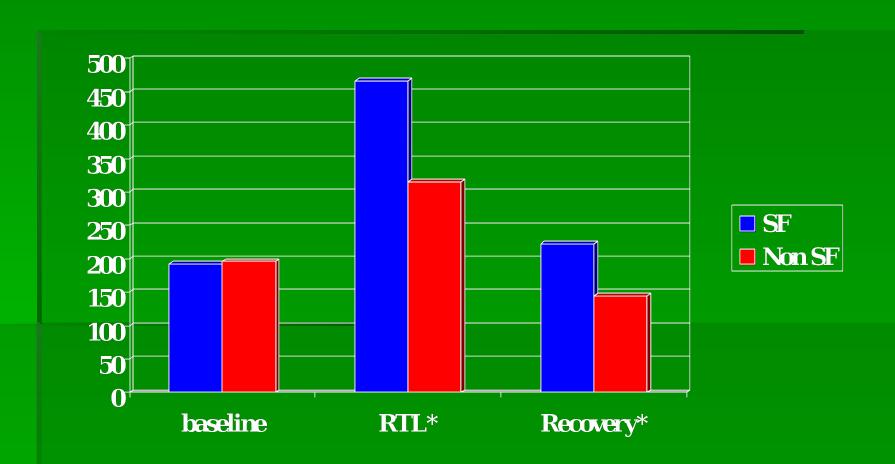
rphism predicts seep structure and waking performance.

besearch centre. School of Biomedical and Mol cularistiences. University of Eure 1, Guijoford, Limitative and sleep has beestasis interact to regulate sleep—wake tydes (1-4), but the genetic basis of prences of percent at the contribution of the

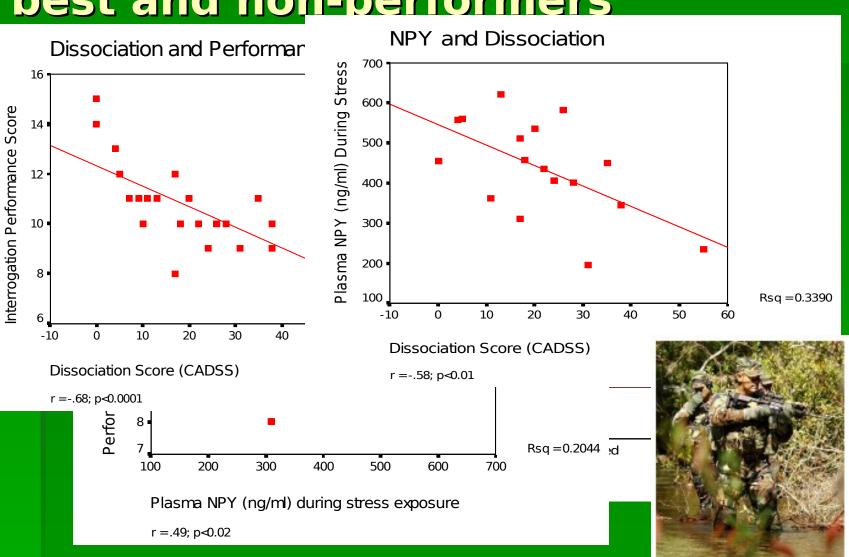
a Korean Asian than in a
 Swedish Caucasian
 population, with 66.7 and 9.3
 % deletion/deletion (del/del)
 homozygotes respectively.

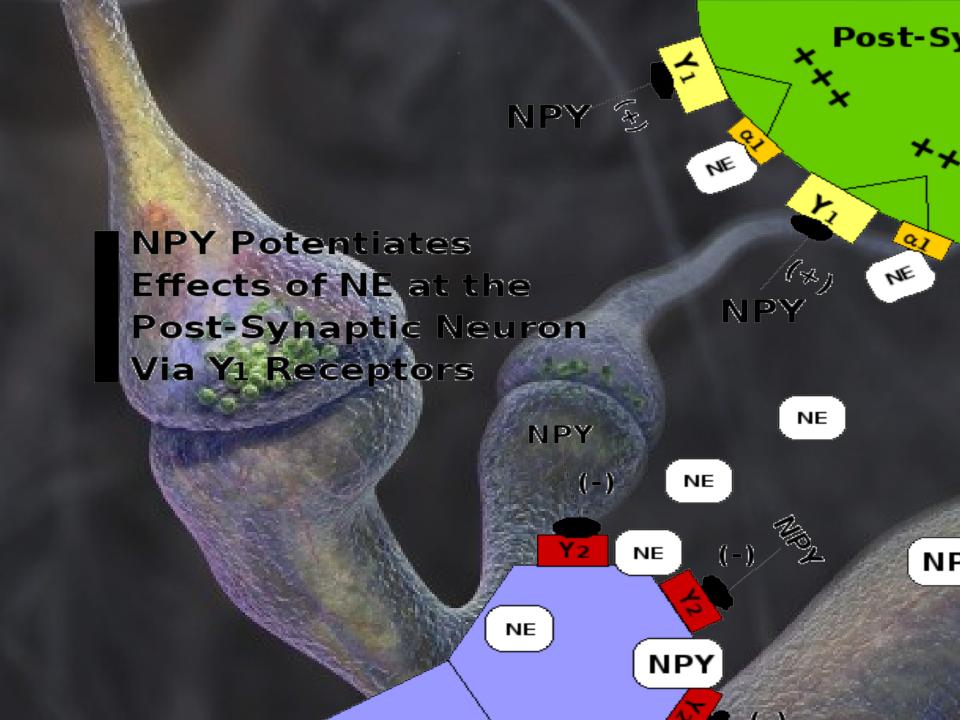
Nutrigenomics: speculation

Neuropeptide Y (Morgan et al., 2000; 2001)



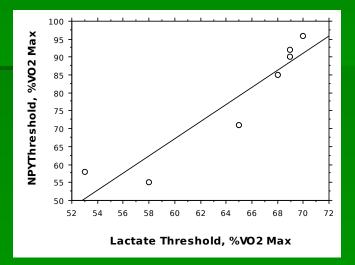
NPY and Human Performance: best and non-performers



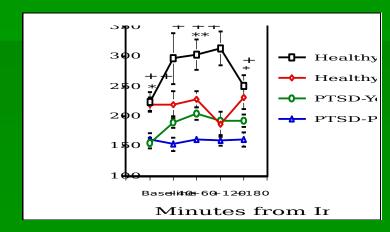


Clinical NPY Data

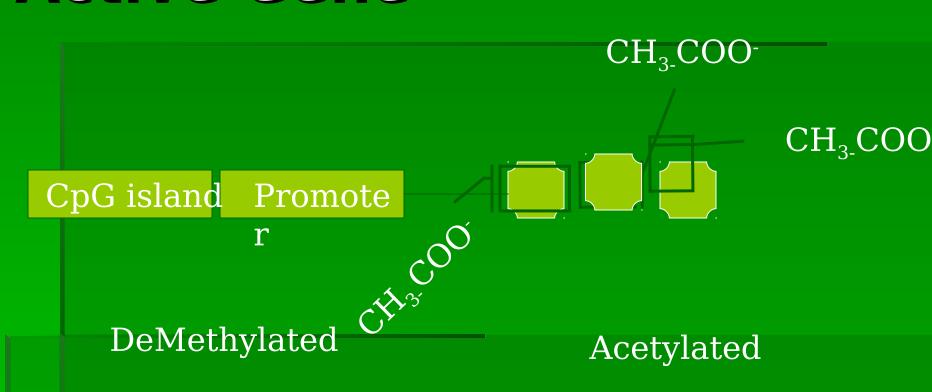
- Increased NPY found in response to 75% VO2 max exercise (Lundberg 1985), the cold pressor test (Morris 1986), and in response to noradrenergic activation by alpha-2 receptor antagonist yohimbine (Rasmusson 1998)
- Reduced NPY noted in CSF of patients suffering from major depression (Widerlov 1992), suicide victims (Widdowson 1992) Negative correlation noted between anxiety scores and CSF NPY levels in patients with depression (Heilig 1990)



 Reduced baseline NPY & blunted NPY response to yohimbine stimulation in veterans with PTSD (Rasmusson 2000).

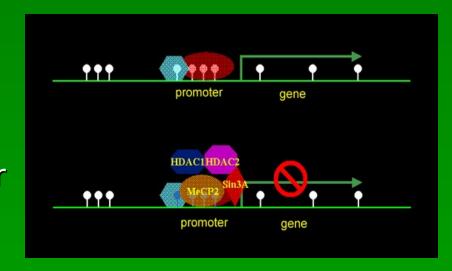


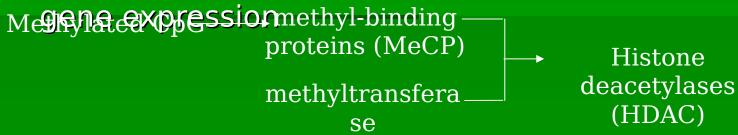
Epigenetic Regulation "Active Gene"



Epigenetics

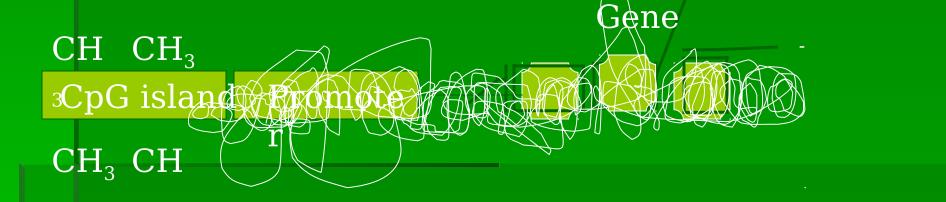
- Promoters are regulatory elements upstream the 5' end of TSS.
- Methylation of promoter CpGs remodels the chromatin structure for





"Pattern Detection and Co-methylation Analysis of Epigenetic Features in Human Embryonic Stem Cells." 2008 Presentation by Ben Niu ☐ Qiang Yang, Jinyan Li, Hong Xue, Simon Chi-keung Shiu, Weichuan Yu, Huiqing Liu, Sankar Kumar Pal. Hong Kong Polytechnic University

Epigenetic Regulation"Moth-balled gene"



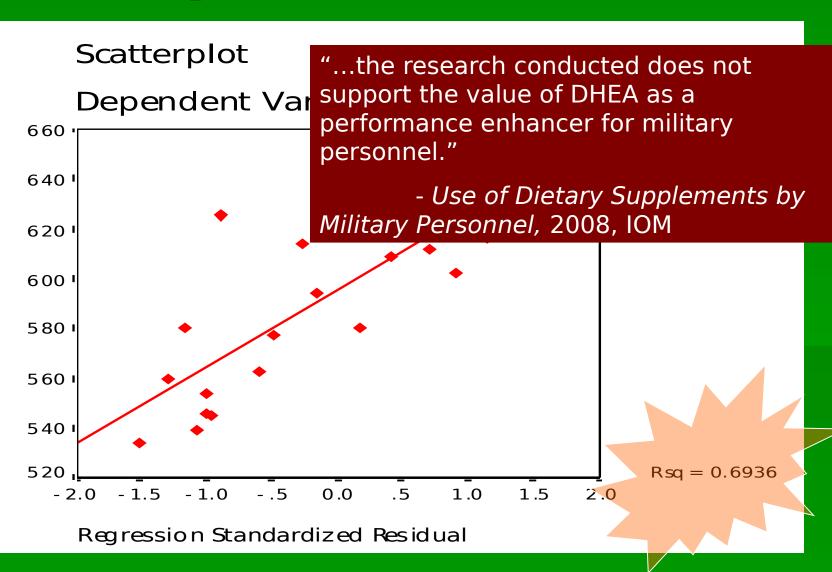
Methylated

DeAcetylat ed

Unmethylated

Diet-based deacetylase inhibitors like bacterial fermentation of carbohydrate within the gastrointestinal tract, ketones (butyrate, valproic acid), ketogenic diet?

A side point...



Nutrigenetics: yet more speculation

The prevalence of folateremedial MTHFR enzyme variants in humans.

*Marini N.J., Gin J, Ziegle J, Keho KH, David Ginzinger D., Gilbert D. and Rine J. *PNAS*, v.105(2): June 10, 2008

- Sampled 11
 methylenetetrahydrofolate
 reductase (MTHFR) SNPs, from
 564 individuals of diverse
 ethnicities
- "Multiple less-frequent alleles, in aggregate, might significantly contribute to metabolic dysfunction. Furthermore, vitamin remediation of mutant enzymes may be a common phenomenon in certain domains of proteins."

 Broccoli Consumption Interacts with GSTM1 to Perturb Oncogenic Signalling Pathways in the Prostate.

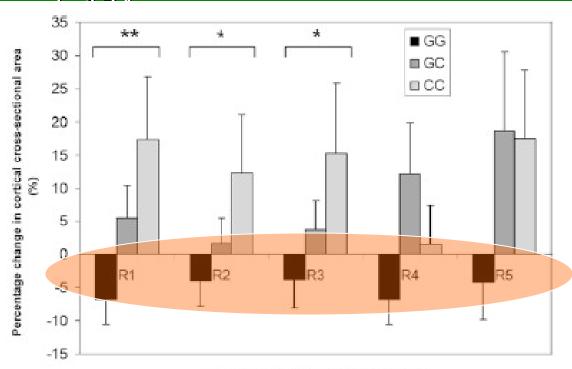
Traka M, Gasper AV, Melchini A, Bacon JR, Needs PW, et al. (2008) *PLoS ONE* 3(7): e2568.

Genetic Variant in the Glucose Transporter Type 2 (GLUT 2) is Associated with Higher Intakes of Sugars in Two Distinct Populations.

Ahmed El-Sohemy, A., Eny, K.M., Wolever, T., and Fontaine-Bisson, B. (2008) *Physiological Genomics* May 2008

IL-6 SNP and bone density

Fig. 1 Mean percentage change in right femoral cortical cross-sectional area in five slices taken at 50 mm intervals (RIproximal to R5distal) associated with a 10-week strenuous exercise training programme in 130 young healthy males genotyped for the -174 G>C IL-6 gene promoter polymorphism. **P = 0.027 by ANOVA;P = 0.007 for linear trend. *P<0.05 for linear trend; ANOVA NS

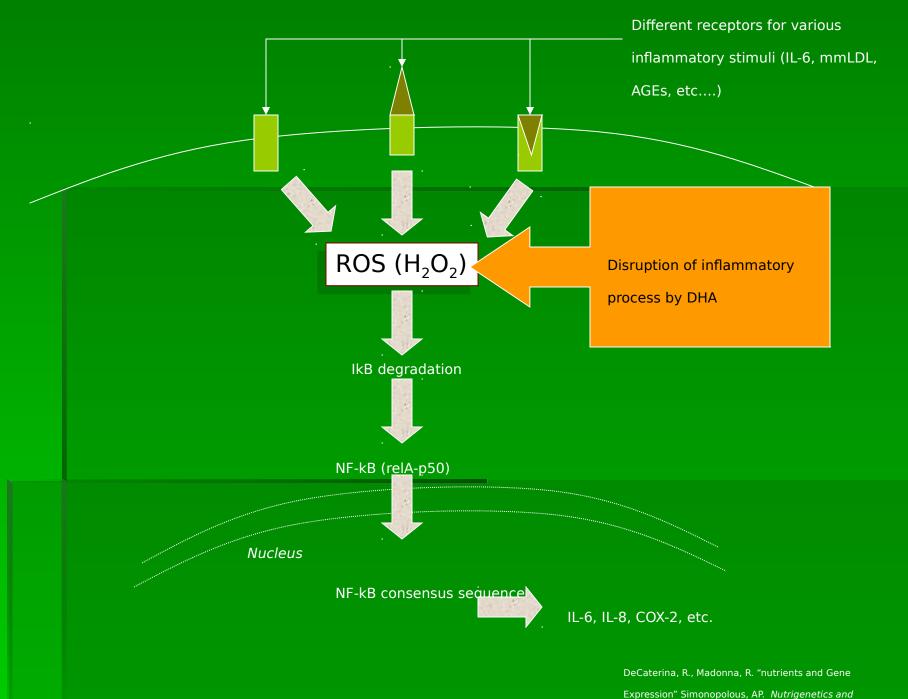


Right Femoral Cortical Slice Number

weinberg equilibriani

GG: 36% GC: 47%

CC: 22.17%



expression simonopolous, AP. Natingenetics and

Paradigm Shift

- As nutrigenomics is inherently PROactive, the challenge is going to be getting from:
 - Description: this creature eats x, then y (sometimes) happens...or not
 - Prediction: therefore, if creature eats x, then y will/will not have a better chance of happening
 - Presc creat pt want y,

Beyond speculation





Traveling out of town?

>> Learn

Update your journal

Resisted temptation today?
Feeling energized?

Struggling to find motivation?

Looking for a spa nearby?

Heed weight loss suggestions? Tricks for keeping stress away? Motivation boosts? Ready to make your cell phone

just text:

"well" to 878787

Out at your favorite restaurant?
Want nutrition info on a menu item?
Looking for a healthier alternative?
» Learn more

The best way to predict the future is to create it.

Abraham Lincoln

Oh, yeah...

Practical Issues

- Novel nutrient-gene interactions
- New diagnostic tests for responses to diets and new biomarkers like mRNA for stress (oxidative or otherwise) that can be used to detect biopotency/applicability
- Identifying specific populations with special needs
- Improving the definitions and methodologies related to dietary assessment
- Providing information to make "food +"

Ethical Issues

- Consent
- •GINA may mean something different to military (i.e. 2001 Burlington Northern Santa Fe lawsuit where the company tested workers with carpal tunnel syndrome for genetic predispositiong; athletes prevented from competing if they have hypertrophic cardiomyopathy (HCM) Chicago Bulls and Eddy Curry.
- Confidentiality
- Solidarity (same species, different creatures)
- •Knowing what's coming (AD risk for example)
- •Access: right now, the rich can afford this and may drive it forward but if there's really a there there, shouldn't it be made available to those who need it most: the disadvantaged, the diseased, and the defenders?

Methodological Issues

- Study design limitations
- Need to incorporate epigenetics
- SNP identification and haplotyping
- Dietary intake assessment
- Better biomarkers
- Demonstrate analytical AND clinical validity
- Clinical utility

Thanks

Thanks due to Ann Rasmussen (Yale), Gary Hazlett (Cody Woodward LLC), and Andy Morgan (Yale)

Contact information: Adam Russell, Scitor Corporation arussell@scitor.com, 202-316-5088